

Figure 1

Mouse Human	MGGCEVREFLLQFGFFLPLLTAWTGDCSHVSN-QVVLLDTTTVMGELGWKTYPLNGWDAI MGGCEVREFLLQFGFFLPLLTAWPGDCSHVSNNQVVLLDTTTVLGELGWKTYPLNGWDAI ************************************	59 60
Mouse Human	TEMDEHNRPIHTYQVCNVMEPNQNNWLRTNWISRDAAQKIYVEMKFTLRDCNSIPWVLGT TEMDEHNRPIHTYQVCNVMEPNQNNWLRTNWISRDAAQKIYVEMKFTLRDCNSIPWVLGT ************************************	119 120
Mouse Human	CKETFNLYYIESDESHGTKFKPSQYIKIDTIAADESFTQMDLGDRILKLNTEIREVGPIE CKETFNLFYMESDESHGIKFKPNQYTKIDTIAADESFTQMDLGDRILKLNTEIREVGPIE ******:******************************	179 180
Mouse Human	RKGFYLAFQDIGACIALVSVRVFYKKCPFTVRSLAMFPDTIPRVDSSSLVEVRGSCVKSA RKGFYLAFQDIGACIALVSVRVFYKKCPFTVRNLAMFPDTIPRVDSSSLVEVRGSCVKSA ************************************	239 240
Mouse Human	EERDTPKLYCGADGDWLVPLGRCICSTGYEEIEGSCHACRPGFYKAFAGNTKCSKCPPHS EERDTPKLYCGADGDWLVPLGRCICSTGYEEIEGSCHACRPGFYKAFAGNTKCSKCPPHS ***********************************	299 300
Mouse Human	STYVEATSVCHCEKGYFRAEKDPPSMACTRPPSAPRNVAFNINETALILEWSPPSDTGGR LTYMEATSVCQCEKGYFRAEKDPPSMACTRPPSAPRNVVFNINETALILEWSPPSDTGGR **:*****:***************************	359 360
Mouse Human	KDLTYSVICKKCGLDTTQCEDCGGGLRFIPRHTGLINNSVVVLDFVSHVNYTFEIEAMNG KDLTYSVICKKCGLDTSQCEDCGGGLRFIPRHTGLINNSVIVLDFVSHVNYTFEIEAMNG ************************************	
Mouse Human	VSELSISPKPFTAITVTTDHDAPSLIGMMRKDWASQNSLALSWQAPAFSNGAILDYETKY VSELSFSPKPFTAITVTTDQDAPSLIGVVRKDWASQNSIALSWQAPAFSNGAILDYEIKY ****:********************************	479 480
Mouse Human	YEKEHEQLTYSSTRSKAPSVIVTGLKPATTYIFHIRVRTATGYSGYSQKFEFETGDETSD YEKEHEQLTYSSTRSKAPSVIITGLKPATKYVFHIRVRTATGYSGYSQKFEFETGDETSD ************************************	539 540
Mouse Human	MAAEQGQILVIATAAVGGFTLLVILTLFFLITGRCQWYIKAKMKSEEKRRTHLQNGHLRF MAAEQGQILVIATAAVGGFTLLVILTLFFLITGRCQWYIKAKMKSEEKRRNHLQNGHLRF ************************************	599 600
Mouse Human	PGIKTYIDPDTYEDPSLAVHEFAKEIDPSRIRIERVIGAGEFGEVCSGRLKTPGKREIPV PGIKTYIDPDTYEDPSLAVHEFAKEIDPSRIRIERVIGAGEFGEVCSGRLKTPGKREIPV ************************************	659 660
Mouse Human	AIKTLKGGHMDRQRRDFLREASIMGQFDHPNIIRLEGVVTKRSFPAIGVEAFCPSFLRAG AIKTLKGGHMDRQRRDFLREASIMGQFDHPNIIRLEGVVTKRSFPAIGVEAFCPSFLRAG ************************************	719 720
Mouse Human	FLNGIQAPHPVTAGGSLPPRIPAGRPVMIVVEYMENGSLDSFLRKHDGHFTVIQLVGMLR FLNSIQAPHPVPGGGSLPPRIPAGRPVMIVVEYMENGSLDSFLRKHDGHFTVIQLVGMLR ***.******************************	779 780
Mouse Human	GIASGMKYLSDMGYVHRDLAARNILVNSNLVCKVSDFGLSRVLEDDPEAAYTTTGGKIPI GIASGMKYLSDMGYVHRDLAARNILVNSNLVCKVSDFGLSRVLEDDPEAAYTTTGGKIPI ***********************************	
Mouse Human	RWTAPEAIAYRKFSSASDAWSYGIVMWEVMSYGERPYWEMSNQDVILSIEEGYRLPAPMG RWTAPEAIAYRKFSSASDAWSYGIVMWEVMSYGERPYWEMSNQDVILSIEEGYRLPAPMG ************************************	899 900
Mouse Human	CPPSLQQLMLHCWQKERNHRPKFTDIVSFLDKLIRNPSALHTLVEDILVMPESPGDVPEY CPASLHQLMLHCWQKERNHRPKFTDIVSFLDKLIRNPSALHTLVEDILVMPESPGEVPEY **.**:********************************	
Mouse Human	PLFVTVGDWLDSIKMGQYKSNFMAAGFTTFDLISRMSIDDIRRIGVILIGHQRRIVSSIQ PLFVTVGDWLDSIKMGQYKNNFVAAGFTTFDLISRMSIDDIRRIGVILIGHQRRIVSSIQ ***********************************	1019 1020
Mouse Human	TLRLHMMHIQEKGFHV 1035 TLRLHMMHIQEKGFHV 1036 *********	

## Figure 3

Human: Mouse:	aggacccccatcctacccagaacacctgcctgcgctgccgccacttctctttaagggaga	
Human: Mouse:	ggaaaagagagcctaggagaaccatggggggctgcgaagtccgggaatttcttttgcaat	
Human: Mouse:	ttggtttcttcttgcctctgctgacagcgtggccaggcgactgcagtcacgtctccaaca	
Human: Mouse:	accaagttgtgttgcttgatacaaccactgtactgggagagctaggatggaaaacatatc	
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Human: Mouse:	ttcgtgaggtggggcctatagaaaggaaaggattttatctggcttttcaagacattgggg 	
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Human: Mouse:	acttggccatgtttcctgataccattccaagggttgattcctcctctttggttgaagtac	
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Human: Mouse:	atggagattggctggttcctcttggaaggtgcatctgcagtacaggatatgaagaaattg 	
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	aaacaggagatgaaacttctgacatggcagcagaacaaggacagattctcgtgatagcca   !	

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Mouse:	1939	ctgcagccgtcgggggattcactctttagtcatcctcaccctgttcttcctcatcatcg 1998
Luman	1903	ggagatgtcagtggtacataaaagccaagatgaagtcagaagagaagaagaaaaccact 1862
Mouse.	1999	ggaggtgtcaatggtacataaaggccaaaatgaagccagaagaagaagaagaacccass
Human:	1863	tacagaatgggcatttgcgcttcccgggaattaaaacttacattgatccagatacatatg 1922
Mouse:	2059	tacagaacggccacctgcgcttcccgggaattaaaacatacat
Human:	1923	aagacccatccctagcagtccatgaatttgcaaaggagattgatccctcaagaattcgta 1982
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Human:	1983	ttgagagagtcattggggcaggtgaatttggagaagtctgtagtgggcgtttgaagacac 2042
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Human:	2043	cagggaaaagagatcccagttgccattaaaactttgaaaggtggccacatggatcggc 2102
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Human•	2343	atatggagaatggatccctagactcctttttgcggaagcatgatggccacttcacagtca 2402
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		aagtttctgattttggtctctccagagtgctggaagatgatccagaagctgcttatacaa 2582
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Mouse:	2779	caacgggagggaaaatccctatacggtggacagccccagaagctattgcttacagaaagt	2838
Human:	2643	tetectcagcaagcgatgcatggagctatggcattgtcatgtgggaggtcatgtcctatg	2702
Mouse:	2839		2898
Human:	2703	gagagagaccttattgggaaatgtctaaccaagatgtcattctgtccattgaagaagggt	2762
Mouse:	2899		2958
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Mouse:	3019	ggcagaaggagagaaaccacaggccaaaattcactgacatcgtcagcttccttgacaaac	3078
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Mouse:	3139	cccctggtgatgttcctgaatatccattgtttgtcacagtaggtgactggctgg	3198
Human:	3003	taaagatggggcaatacaagaataacttcgtggcagcagggtttacaacatttgacctga	3062
Mouse:	3199	taaagatggggcaatacaagagtaacttcatggcagcgggttttacaacgtttgatctga	3258
Human:	3063	tttcaagaatgagcattgatgacattagaagaattggagtcatacttattggacaccaga	3122
Mouse:	3259	$\verb tttcacga  at gag catcgat gat at tagg cga at tgg agt cattct cattgg a cat caga$	3318
Human:	3123	gacgaatagtcagcagcatacagactttacgtttacacatgatgcacatacaggagaagg	3182
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Human:	3183	gatttcatgtatgaaagtaccacaagcacctgtgttttgtgcctcagcatttctaaaatg	3242
Mouse:	3379	gatttcatgtatgaaagtactataaacacctgtatcttgtgcctcagcatttctacaaag	3438
		aacgatatcctctctactactctctcttctgattctccaaacatcacttcacaaactgca	
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Human:	3363	atcatgctacataaatccgttctgaataacctgcaactaaaaccctggcccactgcagat	3422
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Human: 3483 gggaagtgttcacggacttaacctaaaaaaatttatcc--aggtggggcttccttagtga 3540

Mouse: 3678 gggaagtgttcatggacttcacctaaaggaatttgtccttaagtgtggcatcttcagaga 3737

Human: 3601 gatcaagtagct 3612 1111 1111111

Mouse: 3798 gatcgagtagct 3809

## Figure 4

KSE132 XM_035574	MGGCEVREFLLQFGFFLPLLTAWPGDCSHVSNNQVVLLDTTTVLGELGWKTYPLNGWDAI	60
KSE132 XM_035574	TEMDEHNRPIHTYQVCNVMEPNQNNWLRTNWISRDAAQKIYVEMKFTLRDCNSIPWVLGT	120
KSE132 XM 035574	CKETFNLFYMESDESHGIKFKPNQYTKIDTIAADESFTQMDLGDRILKLNTEIREVGPIE	180
KSE132 XM_035574	RKGFYLAFQDIGACIALVSVRVFYKKCPFTVRNLAMFPDTIPRVDSSSLVEVRGSCVKSA	240
KSE132 XM_035574	EERDTPKLYCGADGDWLVPLGRCICSTGYEEIEGSCHACRPGFYKAFAGNTKCSKCPPHS	300
KSE132 XM_035574	LTYMEATSVCQCEKGYFRAEKDPPSMACTRPPSAPRNVVFNINETALILEWSPPSDTGGR	360
KSE132 XM_035574	KDLTYSVICKKCGLDTSQCEDCGGGLRFIPRHTGLINNSVIVLDFVSHVNYTFEIEAMNG	420
KSE132 XM_035574	VSELSFSPKPFTAITVTTDQDAPSLIGVVRKDWASQNSIALSWQAPAFSNGAILDYEIKY	480
KSE132 XM_035574	YEKEHEQLTYSSTRSKAPSVIITGLKPATKYVFHIRVRTATGYSGYSQKFEFETGDETSD	540 26
KSE132 XM_035574	MAAEQGQILVIATAAVGGFTLLVILTLFFLITGRCQWYIKAKMKSEEKRRNHLQNGHLRF MAAEQGQILVIATAAVGGFTLLVILTLFFLITGRCQWYIKAKMKSEEKRRNHLQNGHLRF ************************************	600 86
kse132 xm_035574	PGIKTYIDPDTYEDPSLAVHEFAKEIDPSRIRIERVIGAGEFGEVCSGRLKTPGKREIPV PGIKTYIDPDTYEDPSLAVHEFAKEIDPSRIRIERVIGAGEFGEVCSGRLKTPGKREIPV ************************************	660 146
KSE132 XM_035574	AIKTLKGGHMDRQRRDFLREASIMGQFDHPNIIRLEGVVTKRSFPAIGVEAFCPSFLRAG AIKTLKGGHMDRQRRDFLREASIMGQFDHPNIIRLEGVVTKRSFPAIGVEAFCPSFLRAG ************************************	720 206
KSE132 XM_035574	FLNSIQAPHPVPGGGSLPPRIPAGRPVMIVVEYMENGSLDSFLRKHDGHFTVIQLVGMLR FLNSIQAPHPVPGGGSLPPRIPAGRPVMIVVEYMENGSLDSFLRKHDGHFTVIQLVGMLR ************************************	780 266
KSE132 XM_035574	GIASGMKYLSDMGYVHRDLAARNILVNSNLVCKVSDFGLSRVLEDDPEAAYTTTGGKIPI GIASGMKYLSDMGYVHRDLAARNILVNSNLVCKVSDFGLSRVLEDDPEAAYTTTD	840 321
KSE132 XM_035574	RWTAPEAIAYRKFSSASDAWSYGIVMWEVMSYGERPYWEMSNQDVILSIEEGYRLPAPMG	900
KSE132 XM_035574	CPASLHQLMLHCWQKERNHRPKFTDIVSFLDKLIRNPSALHTLVEDILVMPESPGEVPEY	960
KSE132 XM_035574	PLFVTVGDWLDSIKMGQYKNNFVAAGFTTFDLISRMSIDDIRRIGVILIGHQRRIVSSIÇ	1020
KSE132 XM_035574	TLRLHMMHIQEKGFHV 1036	